

GenCore version 4.5  
 Copyright (c) 1993 - 2000 Compugen Ltd.

run on: February 13, 2002, 10:08:02 ; Search time 12.49 Seconds  
 (without alignments)  
 34.232 Million cell updates/sec

title: US-09-486-094-12  
 perfect score: 51  
 sequence: 1 XCXXXXCXXXXCXXXXCXX 19

scoring table: BLOSUM62  
 Gapop 10.0 , Gapext 0.5

scoring table: 212252 seqs, 22503292 residues

total number of hits satisfying chosen parameters: 212252

int minimum DB seq length: 0  
 maximum DB seq length: 200000000

post-processing: Minimum Match 0%  
 Maximum Match 100%  
 Listing first 45 summaries

database : Issued Patents AA:\*

- 1: /cgp2\_6/ptodata/2/iaa/5A.COMB.pep:\*
- 2: /cgp2\_6/ptodata/2/iaa/5B.COMB.pep:\*
- 3: /cgp2\_6/ptodata/2/iaa/6A.COMB.pep:\*
- 4: /cgp2\_6/ptodata/2/iaa/6B.COMB.pep:\*
- 5: /cgp2\_6/ptodata/2/iaa/PCTMUS.COMB.pep:\*
- 6: /cgp2\_6/ptodata/2/iaa/backfiles1.pep:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

result No.	Score	Query Match	Length	DB ID	Description
1	30	58.8	163	2 US-08-727-688-25	Sequence 25, Appl
2	28	54.9	1345	2 US-08-727-688-25	Sequence 3, Appl
3	27	52.9	45	1 US-08-451-947-97	Sequence 97, Appl
4	27	52.9	45	2 US-08-424-826A-97	Sequence 97, Appl
5	27	52.9	45	3 US-08-424-826A-97	Sequence 97, Appl
6	27	52.9	45	5 PCT-US91-06350-97	Sequence 91, Appl
7	27	52.9	47	3 US-08-482-858B-91	Sequence 11, Appl
8	27	52.9	341	2 US-08-209-521-11	Sequence 6, Appl
9	27	52.9	801	1 US-07-906-349A-6	Sequence 52, Appl
10	27	52.9	4544	1 US-08-459-686-52	Sequence 52, Appl
11	27	52.9	4544	2 US-08-459-686-52	Sequence 5, Appl
12	26	51.0	29	4 US-09-136-759A-5	Sequence 16, Appl
13	26	51.0	29	4 US-09-136-759A-16	Sequence 40, Appl
14	26	51.0	39	1 US-08-050-319B-40	Sequence 40, Appl
15	26	51.0	39	2 US-08-465-982-40	Sequence 112, Appl
16	26	51.0	143	4 US-08-990-893-112	Sequence 52, Appl
17	26	51.0	153	1 US-08-050-319B-52	Sequence 52, Appl
18	26	51.0	153	2 US-08-455-982-52	Sequence 4, Appl
19	26	51.0	153	2 US-08-219-237B-40	Sequence 12, Appl
20	26	51.0	153	4 US-08-477-347-12	Sequence 3, Appl
21	26	51.0	153	4 US-08-08-459-863-3	Sequence 4, Appl
22	26	51.0	153	4 US-08-458-560C-4	Sequence 50, Appl
23	26	51.0	157	1 US-08-050-319B-50	Sequence 50, Appl
24	26	51.0	157	2 US-08-455-982-54	Sequence 50, Appl
25	26	51.0	158	1 US-08-050-319B-54	Sequence 54, Appl
26	26	51.0	161	4 US-08-455-982-54	Sequence 2, Appl

Db 72 C5555C5WP7SCWSTC 88

RESULT 2

US-08-977-767-3 Application US/089777767

Patent No. 5,975,684

GENERAL INFORMATION:

APPLICANT: Bandman, Olga

APPLICANT: Yee, Henry

APPLICANT: Greenwald, Sara

APPLICANT: Corley, Neil C.

TITLE OF INVENTION: CARBONIC ANHYDRASE VIII

NUMBER OF SEQUENCES: 3

CORRESPONDENCE ADDRESS:

ADDRESSEE: Incyte Pharmaceuticals, Inc.

STREET: 3174 Porter Drive

CITY: Palo Alto

STATE: CA

COUNTRY: USA

ZIP: 94304

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER: IBM Compatible

OPERATING SYSTEM: DOS

SOFTWARE: FASTSEQ for Windows Version 2.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/977-767

FILING DATE: Herewith

CLASSIFICATION: 424

PRIOR APPLICATION DATA:

APPLICATION NUMBER:

FILING DATE:

ATTORNEY/AGENT INFORMATION:

NAME: Billings, Lucy J.

REGISTRATION NUMBER: 36,749

REFERENCE/DOCKET NUMBER: PF-0423 US

TELECOMMUNICATION INFORMATION:

TELEPHONE: 650-855-0555

TELEFAX: 650-845-4166

TELEX:

INFORMATION FOR SEQ ID NO: 3:

SEQUENCE CHARACTERISTICS:

LENGTH: 1345 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

IMMEDIATE SOURCE:

LIBRARY: Genbank

CLONE: 1532042

US-08-977-767-3

RESULT 3

Query Match 54.9%; Score 28; DB 2; Length 1345;

Best Local Similarity 23.5%; Pred. No. 57;

Matches 4; Conservative 0; Mismatches 13; Indels 0; Gaps 0;

Qy 2 CXXXXCXXXXCXXXXC 18

Db 768 CAATGGCTGGCCATC 784

RESULT 4

US-08-451-947-97

Sequence 97, Application US/08451947

Patent No. 5,702,906

GENERAL INFORMATION:

APPLICANT: GENENTECH, INC.

APPLICANT: ROSENTHAL, ARNON

TITLE OF INVENTION: NOVEL NEUROTROPHIC FACTOR

NUMBER OF SEQUENCES: 100

CORRESPONDENCE ADDRESS:

ADDRESSEE: Genentech, Inc.

STREET: 460 Point San Bruno Blvd

CITY: South San Francisco

STATE: California

COUNTRY: USA

ZIP: 94080

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:



Matches 4; Conservative 0; Mismatches 13; Indels 0; Gaps 0; Db 24 CATGACCATCTCGAGC 40

RESULT 8  
US-08-209-521-11  
Sequence 11, Application US/08209521  
; Patent No. 5922855

GENERAL INFORMATION:  
;  
; APPLICANT: Liskay, Robert M.  
; APPLICANT: Bronner, C. Eric  
; APPLICANT: Baker, Sean M.  
; APPLICANT: Bollag, Roni J.  
; APPLICANT: Kolodner, Richard D.  
; TITLE OF INVENTION: MAMMALIAN DNA MISMATCH REPAIR GENES  
;  
; NUMBER OF SEQUENCES: 30  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Kolisch, Hartwell, Dickinson, McCormack &  
; STREET: 520 S.W. Yamhill, Suite 200  
; CITY: Portland  
; STATE: Oregon  
; COUNTRY: US  
; ZIP: 97204  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent Release #1.0, version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/209, 521  
; FILING DATE: 08-MAR-1994  
; CLASSIFICATION: 435  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Van Ryselbergh, Pierre C.  
; REGISTRATION NUMBER: 33,557  
; REFERENCE/DOCKET NUMBER: OHSU 306A  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (503) 224-6655  
; TELEFAX: (503) 295-5679  
; INFORMATION FOR SEQ ID NO: 11:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 341 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: cDNA  
; POSITION IN GENOME:  
; MAP POSITION: 3p21.3-23  
; US-08-209-521-11

Query Match 52.9%; Score 27; DB 2; Length 341;  
Best Local Similarity 27.3%; Pred. No. 75;  
Matches 3; Conservative 0; Mismatches 8; Indels 0; Gaps 0;

RESULT 9  
US-07-906-349A-6  
Sequence 6, Application US/07906349A  
; Patent No. 544064

GENERAL INFORMATION:  
;  
; APPLICANT: Schlessinger, Joseph  
; APPLICANT: Skolnik, Edward Y.  
; APPLICANT: Margolis, Benjamin L.  
; TITLE OF INVENTION: A NOVEL EXPRESSION-CLONING METHOD FOR IDENTIFYING TARGET PROTEINS FOR EUKARYOTIC TYROSINE KINASES  
;  
; TITLE OF INVENTION: IDENTIFYING TARGET PROTEINS  
; NUMBER OF SEQUENCES: 16

Query Match 52.9%; Score 27; DB 3; Length 47;  
Best Local Similarity 23.5%; Pred. No. 62;  
Matches 4; Conservative 0; Mismatches 13; Indels 0; Gaps 0; Db 24 CATGACCATCTCGAGC 40

INFORMATION FOR SEQ ID NO: 91:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 47 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; US-08-482-085B-91

INFORMATION FOR SEQ ID NO: 18:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 47 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: peptide  
; US-08-482-085B-91

CORRESPONDENCE ADDRESS:  
 ADDRESSEE: Browdy and Neimark  
 STREET: 419 Seventh Street, N.W.  
 STATE: Washington  
 COUNTRY: USA  
 ZIP: 20004

COMPUTER READABLE FORM:  
 MEDIUM TYPE: Floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: PatentIn Release #1.0, Version #1.25  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/07/906,349A  
 FILING DATE: 30-JUN-1992  
 CLASSIFICATION: 435  
 PRIOR APPLICATION DATA:  
 APPLICATION NUMBER: 07/643,237  
 FILING DATE: 18-JAN-1991  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: 202-628-5197  
 TELEFAX: 202-737-3328  
 INFORMATION FOR SEQ ID NO: 6:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 801 amino acids  
 TYPE: amino acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: protein  
 US-07-906-349A-6

RESULT 10  
 US-08-459-486-52  
 ; Sequence 52, Application US/08469486  
 ; Patent No. 573281  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Thoegersen, Hans Christian  
 ; APPLICANT: Holteit, Thor Las  
 ; TITLE OF INVENTION: Improved method for the refolding of  
 ; TITLE OF INVENTION: Proteins  
 ; NUMBER OF SEQUENCES: 58  
 ; CORRESPONDENCE ADDRESS:  
 ; STREET: 225 Franklin Street  
 ; CITY: Boston  
 ; STATE: Massachusetts  
 ; COUNTRY: USA  
 ; ZIP: 02110-2804  
 ; COMPUTER READABLE FORM:  
 ; MEDIUM TYPE: Floppy disk  
 ; COMPUTER: IBM PC compatible  
 ; OPERATING SYSTEM: PC-DOS/MS-DOS  
 ; SOFTWARE: PatentIn Release #1.0, Version 1  
 ; CURRENT APPLICATION DATA:  
 ; APPLICATION NUMBER: US/08/469 658  
 ; FILING DATE: June 5, 1995  
 ; CLASSIFICATION: 530  
 ; PRIOR APPLICATION DATA:  
 ; APPLICATION NUMBER: 08/192,060  
 ; FILING DATE: February 4, 1994  
 ; CLASSIFICATION: 530  
 ; ATTORNEY/AGENT INFORMATION:  
 ; NAME: Paul T. Clark  
 ; REGISTRATION NUMBER: 30-162  
 ; TELEPHONE: 617 542 8906  
 ; TELEFAX: 200154  
 ; INFORMATION FOR SEQ ID NO: 52:  
 ; SEQUENCE CHARACTERISTICS:  
 ; LENGTH: 4544 amino acids  
 ; TYPE: amino acid  
 ; STRANDEDNESS:  
 ; TOPOLOGY: linear  
 ; MOLECULE TYPE: protein

Query Match 52.9%; Score 27; DB 1; Length 801;  
 Best Local Similarity 23.5%; Pred. No. 81;  
 Matches 4; Conservative 0; Mismatches 13; Indels 0;  
 Gaps 0;

QY 2 CXXXXXXCXXXXCXXXXC 18  
 Db 233 CACAAGCTGGCTCTGAC 249

RESULT 11  
 US-08-469-658-52  
 ; Sequence 52, Application US/08469658  
 ; Patent No. 5917018  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Thoegersen, Hans Christian  
 ; APPLICANT: Holteit, Thor Las  
 ; TITLE OF INVENTION: IMPROVED METHOD FOR THE REFOLDING OF  
 ; TITLE OF INVENTION: PROTEINS  
 ; NUMBER OF SEQUENCES: 58  
 ; CORRESPONDENCE ADDRESS:  
 ; STREET: 225 Franklin Street  
 ; CITY: Boston  
 ; STATE: Massachusetts  
 ; COUNTRY: USA  
 ; ZIP: 02110-2804  
 ; COMPUTER READABLE FORM:  
 ; MEDIUM TYPE: Floppy disk  
 ; COMPUTER: IBM PC compatible  
 ; OPERATING SYSTEM: PC-DOS/MS-DOS  
 ; SOFTWARE: PatentIn Release #1.0, Version 1  
 ; CURRENT APPLICATION DATA:  
 ; APPLICATION NUMBER: US/08/469 658  
 ; FILING DATE: June 5, 1995  
 ; CLASSIFICATION: 530  
 ; PRIOR APPLICATION DATA:  
 ; APPLICATION NUMBER: 08/192,060  
 ; FILING DATE: February 4, 1994  
 ; CLASSIFICATION: 530  
 ; ATTORNEY/AGENT INFORMATION:  
 ; NAME: Paul T. Clark  
 ; REGISTRATION NUMBER: 30-162  
 ; TELEPHONE: 617 542 8906  
 ; TELEFAX: 200154  
 ; INFORMATION FOR SEQ ID NO: 52:  
 ; SEQUENCE CHARACTERISTICS:  
 ; LENGTH: 4544 amino acids  
 ; TYPE: amino acid  
 ; STRANDEDNESS:  
 ; TOPOLOGY: linear  
 ; MOLECULE TYPE: protein

Query Match 52.9%; Score 27; DB 1; Length 4544;  
 Best Local Similarity 23.5%; Pred. No. 96;  
 Matches 4; Conservative 0; Mismatches 13; Indels 0;  
 Gaps 0;

QY 2 CXXXXXXCXXXXCXXXXC 18  
 Db 2980 CADVDECSTTFCPSQRC 2996

US-08-469-658-52

Query Match Score 27; DB 2; Length 4544;  
Best Local Similarity 23.5%; Pred. No. 96;  
Matches 4; Conservative 0; Mismatches 13; Indels 0; Gaps 0;  
SEQ ID NO: 2 CXXXXCXXXXCXXC 18  
DB ID NO: 2980 CADVDECSTTEPCSQRC 2996

RESULT 12  
US-09-136-769A-5  
; Sequence 5, Application US/09136769A  
; Patent No. 6307014  
; GENERAL INFORMATION:  
; APPLICANT: Furie, Bruce  
; APPLICANT: Furie, Barbara  
; APPLICANT: Stenflo, Johan  
; APPLICANT: Rigby, Alan C.  
; APPLICANT: Roepstorff, Peter  
; TITLE OF INVENTION: CONOPEPTIDES  
; FILE REFERENCE: 50065/002001  
; CURRENT APPLICATION NUMBER: US/09/136,769A  
; CURRENT FILING DATE: 1998-08-19  
; NUMBER OF SEQ ID NOS: 26  
; SOFTWARE: FastSEQ for Windows Version 4.0  
; SEQ ID NO 5  
; LENGTH: 29  
; TYPE: PRT  
; ORGANISM: Conus textile  
; FEATURE:  
; NAME/KEY: VARIANT  
; LOCATION: (4)...(4)  
; OTHER INFORMATION: xaa is gamma-carboxyglutamic acid

US-09-136-769A-5

Query Match Score 26; DB 4; Length 29;  
Best Local Similarity 23.5%; Pred. No. 89;  
Matches 4; Conservative 0; Mismatches 13; Indels 0; Gaps 0;  
SEQ ID NO: 2 CXXXXCXXXXCXXC 18  
DB ID NO: 8 CSSSGSCCHKSCCRWT 24

RESULT 13  
US-09-136-769A-16  
; Sequence 16, Application US/09136769A  
; Patent No. 6307014  
; GENERAL INFORMATION:  
; APPLICANT: Furie, Bruce  
; APPLICANT: Furie, Barbara  
; APPLICANT: Stenflo, Johan  
; APPLICANT: Rigby, Alan C.  
; APPLICANT: Roepstorff, Peter  
; TITLE OF INVENTION: CONOPEPTIDES  
; FILE REFERENCE: 50065/002001  
; CURRENT APPLICATION NUMBER: US/09/136,769A  
; CURRENT FILING DATE: 1998-08-19  
; NUMBER OF SEQ ID NOS: 26  
; SOFTWARE: FastSEQ for Windows Version 4.0  
; SEQ ID NO 16  
; LENGTH: 29  
; TYPE: PRT  
; ORGANISM: Conus textile  
; FEATURE:  
; NAME/KEY: VARIANT  
; LOCATION: (4)...(4)  
; OTHER INFORMATION: xaa is gamma-carboxyglutamic acid

US-09-136-769A-16

Query Match Score 26; DB 4; Length 29;  
Best Local Similarity 23.5%; Pred. No. 89;  
Matches 4; Conservative 0; Mismatches 13; Indels 0; Gaps 0;  
SEQ ID NO: 2 CXXXXCXXXXCXXC 18  
DB ID NO: 8 CSSSGSCCHKSCCRWT 24

RESULT 14  
US-08-050-319B-40  
; Sequence 40, Application US/08050319B  
; Patent No. 5633145  
; GENERAL INFORMATION:  
; APPLICANT: M. Feldmann, P. W. Gray,  
; APPLICANT: M.J.C. Turner, F.M. Brennan  
; TITLE OF INVENTION: Modified human TNFalpha (Tumor  
; Necrosis Factor alpha) Receptor  
; NUMBER OF SEQUENCES: 57  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Reed & Robbins  
; STREET: 635 Bryant Street  
; CITY: Palo Alto  
; STATE: California  
; COUNTRY: USA  
; ZIP: 94301  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/050,319B  
; FILING DATE: 10-May-1993  
; CLASSIFICATION: 435  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Robbins, Roberta L.  
; REGISTRATION NUMBER: 33,208  
; REFERENCE/DOCKET NUMBER: 5150-0030  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (415) 617-8999  
; TELEFAX: (415) 327-5231  
; INFORMATION FOR SEQ ID NO: 40:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 39 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: Linear  
; MOLECULE TYPE: protein  
; SEQ ID NO: 40  
; LENGTH: 29  
; TYPE: PRT  
; ORGANISM: Conus textile  
; FEATURE:  
; NAME/KEY: VARIANT  
; LOCATION: (4)...(4)  
; OTHER INFORMATION: xaa is gamma-carboxyglutamic acid

US-08-050-319B-40

RESULT 15  
US-08-465-982-40  
; Sequence 40, Application US/08465982  
; Patent No. 5863786  
; GENERAL INFORMATION:  
; APPLICANT: M. Feldmann, P.W. Gray,  
; APPLICANT: M.J.C. Turner, F.M. Brennan  
; TITLE OF INVENTION: Modified human TNFalpha (Tumor  
; Necrosis Factor alpha) Receptor  
; NUMBER OF SEQUENCES: 57

```

; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Reed & Robbins
; STREET: 635 Bryant Street
; CITY: Palo Alto
; STATE: California
; COUNTRY: USA
; ZIP: 94301

; COMPUTER READABLE FORM:
; MEDIUM TYPE: floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/465,982
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/050,319
; FILING DATE: 10-May-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Robbins, Roberta L.
; REGISTRATION NUMBER: 33,208
; REFERENCE/DOCKET NUMBER: 5150-0030
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 617-8999
; TELEFAX: (415) 327-3231
; INFORMATION FOR SEQ ID NO: 40:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 39 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-465-982-40

* Query Match      51 0%; Score 26; DB 2; Length 39;
* Best Local Similarity 23.5%; Pred. No. 92;
* Matches 4; Conservative 0; Mismatches 13; Indels 0; Gaps 0;
* Qy   2 CXXXXCXXXXCXXXXC 18
*          |           |           |
* Db   13 CVSCSNCKKSLEETKLC 29

```

Search completed: February 13, 2002, 10:09:47  
 Job time: 105 sec

